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Sita

Sita Pappu
Art Unit 1632
3D05/305-5039

of Contact:
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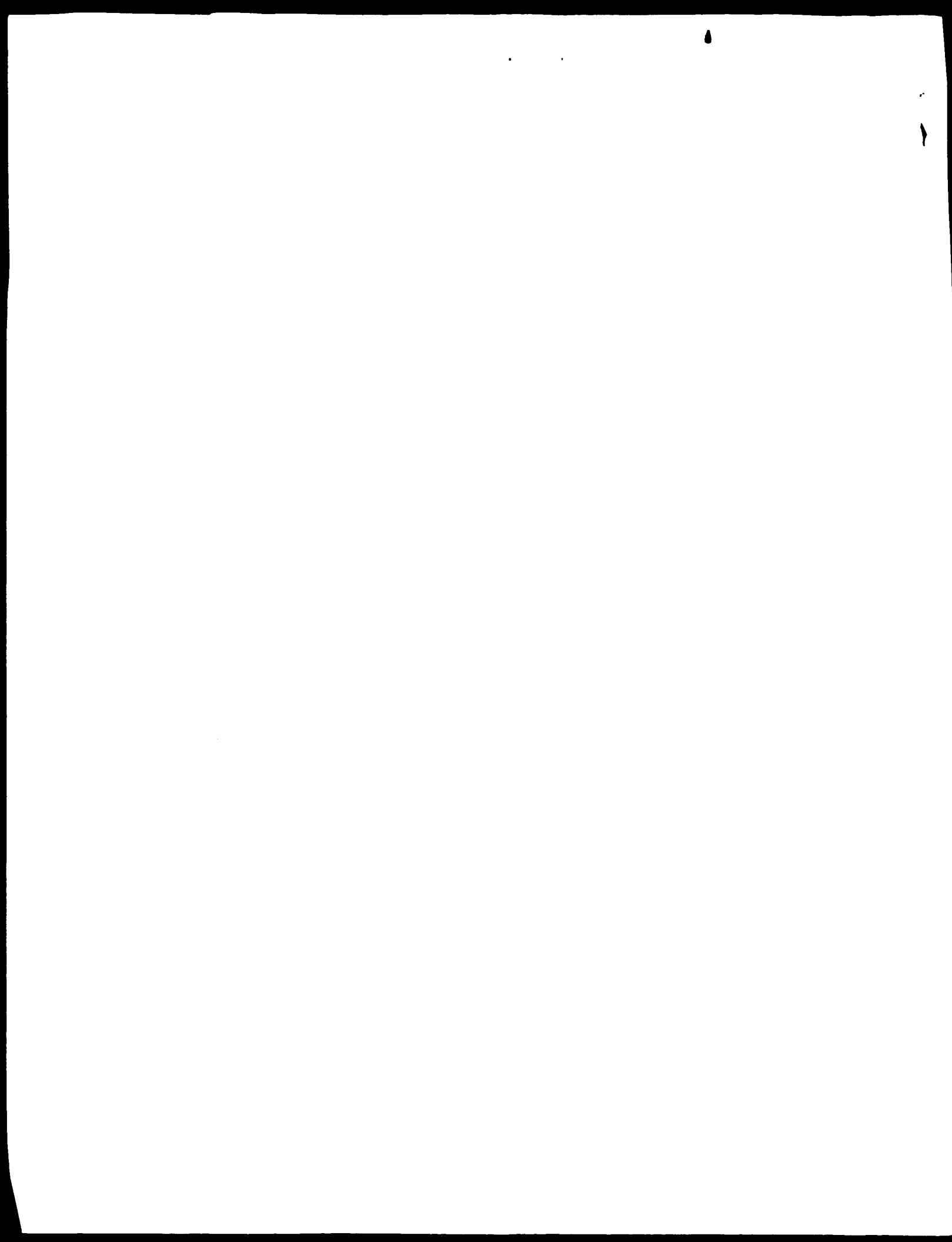
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121 01-JUN-2001 [TIFMHL,et al. 17: 140]

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12 Lactamase (S). Pathram Maludro V. Malsing D:
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13 With 2.7e-42.647e/100
14 N PDB: AAA29006.

XX
15 Product for treatment, preventing and diagnosing cervical cancer
16 represents a nucleotid sequence of molecule which binds to Bcrn No.
17 decreases its intracellular levels or inhibits its activity.

XX
18 Disclosure: Page 64-65: 72pp: English.

XX
19 A Product that binds, causes a decrease in intracellular levels of or
20 inhibits the activity of Bcrn-3a useful for treating, preventing or
21 diagnosis of cervical cancer caused by human papilloma virus (HPV) is
22 claimed. Expression of HPV proteins is generally dependent on the
23 presence of Bcrn-3a in the cell. Methods of identifying Bcrn-3a binding
24 agents or agents which inhibit Bcrn-3a expression are disclosed. Mice mice
25 were infected with Siba cells containing a single integrated HPV6 genome
26 were transfected with a Bcrn-3a antisense construct and after the empty
27 expression vector as control and tumours assessed at regular intervals.
28 Results showed that after 30 days there was no or very little tumour
29 growth in mice transfected with Bcrn-3a antisense construct as compared
30 to the control.

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31 Sequence: 423 AA:

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1. *Staphylococcus aureus* (1000)
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Stability of the Equilibrium

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$\frac{1}{2} \frac{d}{dt} \int_{\mathbb{R}^n} |u|^2 dx = \int_{\mathbb{R}^n} u \Delta u dx = - \int_{\mathbb{R}^n} |\nabla u|^2 dx \leq 0$

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Vector: pAKRΔ1; Site 1: 34000-34020 bp (100%); 24 Mouse BAC Library produced by Center for Genomics

Secondary was obtained in the (TBAAN) catalyzed reaction of the methyl ester using Model butadiene diene as 50/50.

1. *Introduction*

PAVING MATERIAL	1967	1968	1969
Asphalt	114	100	98
Concrete	1	1	1
Gravel	1	1	1
Other	1	1	1
Total	117	103	100

[illegible]

0001-9054/90/0005-0000\$05.00/0

[illegible][illegible]

1. The first step is to identify the key components of the system. This involves understanding the hardware, software, and data involved. For example, in a web application, this might include the server, the database, and the user interface.

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100 101 102 103 104 105 106 107 108 109 110 111 112 113 114 115 116 117 118 119 120 121 122 123 124 125 126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143 144 145 146 147 148 149 150 151 152 153 154 155 156 157 158 159 160 161 162 163 164 165 166 167 168 169 170 171 172 173 174 175 176 177 178 179 180 181 182 183 184 185 186 187 188 189 190 191 192 193 194 195 196 197 198 199 200 201 202 203 204 205 206 207 208 209 210 211 212 213 214 215 216 217 218 219 220 221 222 223 224 225 226 227 228 229 230 231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253 254 255 256 257 258 259 260 261 262 263 264 265 266 267 268 269 270 271 272 273 274 275 276 277 278 279 280 281 282 283 284 285 286 287 288 289 290 291 292 293 294 295 296 297 298 299 300 301 302 303 304 305 306 307 308 309 310 311 312 313 314 315 316 317 318 319 320 321 322 323 324 325 326 327 328 329 330 331 332 333 334 335 336 337 338 339 340 341 342 343 344 345 346 347 348 349 350 351 352 353 354 355 356 357 358 359 360 361 362 363 364 365 366 367 368 369 370 371 372 373 374 375 376 377 378 379 380 381 382 383 384 385 386 387 388 389 390 391 392 393 394 395 396 397 398 399 400 401 402 403 404 405 406 407 408 409 410 411 412 413 414 415 416 417 418 419 420 421 422 423 424 425 426 427 428 429 430 431 432 433 434 435 436 437 438 439 440 441 442 443 444 445 446 447 448 449 450 451 452 453 454 455 456 457 458 459 460 461 462 463 464 465 466 467 468 469 470 471 472 473 474 475 476 477 478 479 480 481 482 483 484 485 486 487 488 489 490 491 492 493 494 495 496 497 498 499 500 501 502 503 504 505 506 507 508 509 510 511 512 513 514 515 516 517 518 519 520 521 522 523 524 525 526 527 528 529 530 531 532 533 534 535 536 537 538 539 540 541 542 543 544 545 546 547 548 549 550 551 552 553 554 555 556 557 558 559 560 561 562 563 564 565 566 567 568 569 570 571 572 573 574 575 576 577 578 579 580 581 582 583 584 585 586 587 588 589 590 591 592 593 594 595 596 597 598 599 600 601 602 603 604 605 606 607 608 609 610 611 612 613 614 615 616 617 618 619 620 621 622 623 624 625 626 627 628 629 630 631 632 633 634 635 636 637 638 639 640 641 642 643 644 645 646 647 648 649 650 651 652 653 654 655 656 657 658 659 660 661 662 663 664 665 666 667 668 669 670 671 672 673 674 675 676 677 678 679 680 681 682 683 684 685 686 687 688 689 690 691 692 693 694 695 696 697 698 699 700 701 702 703 704 705 706 707 708 709 710 711 712 713 714 715 716 717 718 719 720 721 722 723 724 725 726 727 728 729 730 731 732 733 734 735 736 737 738 739 740 741 742 743 744 745 746 747 748 749 750 751 752 753 754 755 756 757 758 759 760 761 762 763 764 765 766 767 768 769 770 771 772 773 774 775 776 777 778 779 780 781 782 783 784 785 786 787 788 789 790 791 792 793 794 795 796 797 798 799 800 801 802 803 804 805 806 807 808 809 810 811 812 813 814 815 816 817 818 819 820 821 822 823 824 825 826 827 828 829 830 831 832 833 834 835 836 837 838 839 840 841 842 843 844 845 846 847 848 849 850 851 852 853 854 855 856 857 858 859 860 861 862 863 864 865 866 867 868 869 870 871 872 873 874 875 876 877 878 879 880 881 882 883 884 885 886 887 888 889 890 891 892 893 894 895 896 897 898 899 900 901 902 903 904 905 906 907 908 909 910 911 912 913 914 915 916 917 918 919 920 921 922 923 924 925 926 927 928 929 930 931 932 933 934 935 936 937 938 939 940 941 942 943 944 945 946 947 948 949 950 951 952 953 954 955 956 957 958 959 960 961 962 963 964 965 966 967 968 969 970 971 972 973 974 975 976 977 978 979 980 981 982 983 984 985 986 987 988 989 990 991 992 993 994 995 996 997 998 999 1000 1001 1002 1003 1004 1005 1006 1007 1008 1009 1010 1011 1012 1013 1014 1015 1016 1017 1018 1019 1020 1021 1022 1023 1024 1025 1026 1027 1028 1029 1030 1031 1032 1033 1034 1035 1036 1037 1038 1039 104

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Received: 1999-09-10; Accepted: 1999-11-11; Published: 2000-01-01

[illegible][illegible]



[illegible]

ANAL. Calcd for $C_{10}H_{10}O_2$: C, 80.0%; H, 8.0%. Found: C, 79.9%; H, 8.1%.

ANAL. Calcd for $C_{10}H_{10}O$: C, 88.10%; H, 7.40%. Found: C, 88.1%; H, 7.4%.

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1. Introduction

XX How often do you witness animal deaths as a result of hunting practices?

[illegible]

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[illegible]

Figure 6

The figure consists of three panels. The top panel is a bar graph showing the relative expression of four pro-inflammatory cytokines: IL-1 β , IL-6, TNF- α , and MCP-1. The y-axis represents relative expression, ranging from 0 to 1.0. The x-axis shows two groups: Control (n=8) and Treatment (n=8). In the Control group, all four cytokines show high relative expression (approximately 0.8-1.0). In the Treatment group, the expression of all four cytokines is significantly reduced (approximately 0.2-0.4).

The middle panel displays Western blot images for the same four cytokines: IL-1 β , IL-6, TNF- α , and MCP-1. Each row corresponds to one cytokine, and each column corresponds to the Control or Treatment group. The bands are more intense in the Control group compared to the Treatment group.

The bottom panel is a bar graph showing the relative expression of three anti-inflammatory cytokines: IL-10, IL-12(p70), and IFN- γ . The y-axis represents relative expression, ranging from 0 to 1.0. The x-axis shows the same two groups: Control (n=8) and Treatment (n=8). In the Control group, IL-10 expression is approximately 0.8, IL-12(p70) is approximately 0.4, and IFN- γ is approximately 0.2. In the Treatment group, IL-10 expression is significantly increased (approximately 0.9), while IL-12(p70) and IFN- γ expression remain relatively low (approximately 0.2-0.3).

Wolcott, L. L., & A. J.

01 - MARK 2005 -

[illegible]

1 / Add 1 row, 1 col: 00000000.

CONFIDENTIAL - SECURITY INFORMATION

Symbol	Meaning	Symbol	Meaning
\mathcal{A}	Algebra	\mathcal{B}	Basis
\mathcal{C}	Category	\mathcal{D}	Diagram
\mathcal{E}	Equation	\mathcal{F}	Field
\mathcal{G}	Group	\mathcal{H}	Homomorphism
\mathcal{I}	Ideal	\mathcal{J}	Jacobian
\mathcal{K}	Kernel	\mathcal{L}	Linear
\mathcal{M}	Module	\mathcal{N}	Norm
\mathcal{O}	Order	\mathcal{P}	Polynomial
\mathcal{Q}	Quotient	\mathcal{R}	Ring
\mathcal{S}	Space	\mathcal{T}	Tensor
\mathcal{U}	Unit	\mathcal{V}	Vector
\mathcal{W}	Weight	\mathcal{X}	Matrix
\mathcal{Y}	Yield	\mathcal{Z}	Zero

[illegible][illegible]

1. The first step is to identify the problem. This involves understanding the current situation and what needs to be changed.

2. The second step is to set goals. These should be specific, measurable, achievable, relevant, and time-bound (SMART).

3. The third step is to develop a plan. This involves identifying the resources needed and the steps to be taken.

4. The fourth step is to implement the plan. This involves putting the plan into action and monitoring progress.

5. The fifth step is to evaluate the results. This involves comparing the actual results with the goals and making adjustments as needed.

$\frac{d}{dt} \left(\frac{\partial L}{\partial \dot{x}} \right) = \frac{\partial L}{\partial x}$

$\frac{1}{2} \times \frac{1}{2}$

[illegible]

in a critical state. The 11,000 soldiers who were ordered to leave the

medical, therapeutic or prophylactic
analysis of physiologic

in the following manner:

[illegible]

2014-15: Mat. Sci. 27, 18; Sci.

[illegible]

$\frac{1}{\sqrt{\pi}}$

[illegible]

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[illegible]

XX New antitumor and/or antiviral food-ids used in treatment of... (p...)

XX vasoconstriction

XX The specificity of desorption and/or... (AAV 269 X9 271)

XX directed... at least 2... selected from target... (AAV 269 X9 271)

XX non... regions of RNAs corresponding to target genes... (AAV 269 X9 271)

XX initiation... regions, including... (AAV 269 X9 271)

XX regions and... and the... (AAV 269 X9 271)

XX of more... conditions or mixtures... (AAV 269 X9 271)

XX... (AAV 269 X9 271)

XX... (AAV 269 X9 271)

XX... (AAV 269 X9 271)

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		Translated from "MAMMAL CELL PROTEIN KINASE TYPE 1A (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN KINASE TYPE 1A) (MAMMAL CELL PROTEIN 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FEATURE	VALUE
ALGORITHM	ALGORITHM 15
LENGTH	2000
DEFINITION	ALGORITHM 15, 10000 bp DNA
REFERENCE	Human Substrains, Chromosome 18, clone RP11-714M3, not CHAI, SEPULCHRE, & undetected PLOYES.
ACCESSION	AF001460
VERSION	AF001460.2, 01/01/2004
KEYWORDS	BLAST, HUMAN, BLAST, HUMAN, ORF, HUMAN, SUBSTRAINS, CHROMOSOME 18, RP11-714M3.
SOURCE	Human Substrains
ORGANISM	Human Substrains

REFERENCE
 (bases 1 to 166144)
 Hattori, M., Ishii, K., Toyoda, A., Taylor, T. B., Hattori, T.,
 Fujiyama, A., Yada, T., Tokioka, Y., Matsumoto, and Shibuya,
 Hemo-saptrons 166,144 genome DNA of 18121
 Published only in database (2000) In press
 2 (bases 1 to 166144)
 Hattori, M., Ishii, K., Toyoda, A., Taylor, T. B., Hattori, T.,
 Fujiyama, A., Yada, T., Tokioka, Y., Matsumoto, and Shibuya, Y.
 In press
 In press
 Submitted (29-MAR-2000) Masahiro Hattori, The Institute of Physical
 and Chemical Research (RIKEN), Genome Sciences Center (GSC),
 Kitasato Univ., 1-15-1 Kitasato, Saitama-shi, Kitasato 426-8601,
 Japan (Email:hattori@gsr.riken.ac.jp,
 hhattori@gsr.riken.ac.jp, Tel: 91-42-778-9924,
 Fax: 91-42-778-9924)
 EMBL
 In May 1999, this sequence version replaced all 1900995.

Project name: RIKEN
Web site: <http://doh.riken.go.jp/>
Contact: hattori@do.riken.go.jp

Project information

Center project name: Humu-att18
Center clone name: R111-71.1M24

Summary statistics

Sequencing vector: PCR products: 100K of total
Chemistry: Dye terminator: 4F dms-slm, 100K of total
Assembly program: Phrap, version 0.799.2.0
Consensus quality: 194875 bases at least 40
Consensus quality: 160311 bases at least 40

Consensus quality: 162445 bases at least Q20
Insert size: 163944; sum of contents
Quality: contigs are sorted in descending order of

NOTE: This is a working draft sequence. It currently consists of 23 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be increased.

Sequence	Updated at	(20-May-2000)
2,27,25	4,55,25	count in out
4,36,27	4,61,14	count in out
6,04,15	7,84,2	count in out
7,94,4	8,80,5	count in out
8,79,66	9,78,0	count in out
9,79,87	10,83,9	count in out
10,94,60	11,55,38	count in out
11,90,59	12,16,17	count in out
11,77,18	12,43,4	count in out
12,85,4	12,81,69	count in out
12,82,70	13,82,79	count in out
13,83,80	13,85,43	count in out
13,83,80	14,77,77	count in out
14,08,78	14,90,44	count in out
14,11,15	14,93,0	count in out
14,94,10	15,40,4	count in out
15,41,55	15,54,57	count in out
15,64,48	15,51,52	count in out
15,92,5	16,08,70	count in out
16,09,71	16,13,45	count in out
16,22,64	15,62,25	count in out
16,67,26	16,61,14	count in out
	14,19	bp in bp

1	22159	22258	gap of	100 bp	in length
2	22559	22658	contig of	21298 bp	in length
3	44527	44626	contig of	41298 bp	in length
4	46627	46726	gap of	100 bp	in length
5	60415	60414	contig of	14688 bp	in length
6	60415	60414	gap of	100 bp	in length
7	69415	72942	contig of	17428 bp	in length
8	72843	72942	gap of	100 bp	in length
9	72943	87865	contig of	9928 bp	in length
10	87866	87965	gap of	100 bp	in length
11	87966	97846	contig of	9871 bp	in length
12	97937	97936	gap of	100 bp	in length
13	103460	103459	contig of	5428 bp	in length
14	103460	103459	gap of	100 bp	in length
15	103460	110558	contig of	7079 bp	in length
16	110559	110658	gap of	100 bp	in length
17	110659	117617	contig of	6939 bp	in length
18	117618	117717	gap of	100 bp	in length
19	117718	123453	contig of	7756 bp	in length
20	123454	123553	gap of	100 bp	in length
21	123554	128169	contig of	4913 bp	in length
22	128170	128269	gap of	100 bp	in length
23	128270	133279	contig of	5010 bp	in length
24	133280	133379	gap of	100 bp	in length
25	133380	137531	contig of	43742 bp	in length
26	137532	137631	gap of	100 bp	in length
27	137632	140377	contig of	31149 bp	in length
28	140378	140377	gap of	100 bp	in length
29	140378	145044	contig of	4167 bp	in length
30	145045	145144	gap of	100 bp	in length

